

SEQUENCE LISTING

<110> Waldman, Scott A.
Park, Jason
Schulz, Stephanie

<120> Compositions And Methods For Identifying And Targeting Cancer Cells Of Alimentary Canal Origin

<130> TJU2389

<150> 60/192,229

<151> 2000-03-27

<160> 2

<170> PatentIn version 3.0

<210> 1

<211> 1745

<212> DNA

<213> Homo sapiens

<400> 1

gcgcccttg cagccttcaa cgtcggtccc caggcagcat ggtgaggtct gctcccggac	60
cctcgccacc atgtacgtga gctacctcct ggacaaggac gtgagcatgt accctagctc	120
cgtgcgccac tctggcgggc tcaacctggc gccgcagaac ttcgtcagcc cccgcagta	180
cccggaactac ggcggttacc acgtggcggc cgcagctgca gcgcagaact tggacagcgc	240
gcagtccccg gggccatcct ggccggcagc gtatggcgcc ccactccggg aggactggaa	300
tggctacgcg cccggaggcg cggccgccgc caacgccgtg gctcacgcgc tcaacggtgg	360
ctccccggcc gcagccatgg gctacagcag cccgcagac taccatccgc accaccacc	420
gcacaccac ccgcaccacc cggccgccgc gccttcctgc gcttctgggc tgctgaaac	480
gctcaacccc ggcctcctg ggcgcgccgc caccgtgcc gccgagcagc tgtctcccg	540
cggccagcgg cggaacctgt gcgagtggat gcggaagccg gcgcagcagt ccctcggcag	600
ccaagtgaac accaggacga aagacaaata tcgagtgggtg tacacggacc accagcggct	660
ggagctggag aaggagtctt actacagtcg ctacatcacc atccggagga aagccgagct	720
agccgccacg ctggggctct ctgagaggca ggtaaaatc tggtttcaga accgcagagc	780
aaaggagagg aaatcaaca agaagaagtt gcagcagcaa cagcagcagc agccaccaca	840
gccgcctccg ccgccaccac agcctcccca gcctcagcca ggtcctctga gaagtgtccc	900
agagcccttg agtccggtgt cttccctgca agcctcagtg tctggctctg tccctggggt	960
tctggggcca actggggggg tgctaaaccc caccgtcacc cagtgacca cgggggtctg	1020
cagcggcaga gcaattccag gctgagccat gaggagcgtg gactctgcta gactcctcag	1080
gagagacccc tcccctccca cccacagcca tagacctaca gacctggctc tcagaggaaa	1140
aatgggagcc aggagtaaga caagtgggat ttggggcctc aagaaatata ctctccaga	1200
tttttacttt ttccatctgg ctttttctgc cactgaggag acagaaagcc tccgctgggc	1260

2389.ST25

ttcattccgg actggcagaa gcattgcctg gactgaccac accaaccagc ttcattctatc 1320
 cgactcttct cttcctagat ctgcaggctg cacctctggc tagagccgag gggagagagg 1380
 gactcaaggg aaaggcaagc ttgaggccaa gatggctgct gcctgctcat ggccctcgga 1440
 ggtccagctg ggccctctgc ctccgggcag caagggtttac actgcggaac gcaaaggcag 1500
 ctaagataga aagctggact gaccaaagac tgcagaaccc ccagggtggc ctgctctttt 1560
 tttctcttcc ctttcccaga ccaggaaagg cttggctggt gtatgcacag ggtgtggtat 1620
 gaggggggtg ttattggact ccaggcctga ccagggggcc cgaacaggac ttgttagaga 1680
 gcctgtcacc agagcttctc tgggctgaat gtatgtcagt gctataaatg ccagagccaa 1740
 cctgg 1745

<210> 2
 <211> 311
 <212> PRT
 <213> Homo sapiens

<400> 2

Met Tyr Val Ser Tyr Leu Leu Asp Lys Asp Val Ser Met Tyr Pro Ser
 1 5 10 15
 Ser Val Arg His Ser Gly Gly Leu Asn Leu Ala Pro Gln Asn Phe Val
 20 25 30
 Ser Pro Pro Gln Tyr Pro Asp Tyr Gly Gly Tyr His Val Ala Ala Ala
 35 40 45
 Ala Ala Ala Gln Asn Leu Asp Ser Ala Gln Ser Pro Gly Pro Ser Trp
 50 55 60
 Pro Ala Ala Tyr Gly Ala Pro Leu Arg Glu Asp Trp Asn Gly Tyr Ala
 65 70 75 80
 Pro Gly Gly Ala Ala Ala Ala Asn Ala Val Ala His Ala Leu Asn Gly
 85 90 95
 Gly Ser Pro Ala Ala Ala Met Gly Tyr Ser Ser Pro Ala Asp Tyr His
 100 105 110
 Pro His His His Pro His His His Pro His His Pro Ala Ala Ala Pro
 115 120 125
 Ser Cys Ala Ser Gly Leu Leu Gln Thr Leu Asn Pro Gly Pro Pro Gly
 130 135 140
 Pro Ala Ala Thr Ala Ala Ala Glu Gln Leu Ser Pro Gly Gly Gln Arg
 145 150 155 160
 Arg Asn Leu Cys Glu Trp Met Arg Lys Pro Ala Gln Gln Ser Leu Gly
 165 170 175
 Ser Gln Val Lys Thr Arg Thr Lys Asp Lys Tyr Arg Val Val Tyr Thr
 180 185 190
 Asp His Gln Arg Leu Glu Leu Glu Lys Glu Phe His Tyr Ser Arg Tyr
 195 200 205

2389.ST25

Ile Thr Ile Arg Arg Lys Ala Glu Leu Ala Ala Thr Leu Gly Leu Ser
210 215 220
Glu Arg Gln Val Lys Ile Trp Phe Gln Asn Arg Arg Ala Lys Glu Arg
225 230 235 240
Lys Ile Asn Lys Lys Lys Leu Gln Gln Gln Gln Gln Gln Pro Pro
245 250 255
Gln Pro Pro Pro Pro Pro Pro Gln Pro Pro Gln Pro Gln Pro Gly Pro
260 265 270
Leu Arg Ser Val Pro Glu Pro Leu Ser Pro Val Ser Ser Leu Gln Ala
275 280 285
Ser Val Ser Gly Ser Val Pro Gly Val Leu Gly Pro Thr Gly Gly Val
290 295 300
Leu Asn Pro Thr Val Thr Gln
305 310

2389.ST25